SEQUENCE LISTING <110> E. I. du Pont de Nemours and Company <120> Membrane-Bound Desaturases <130× BB1264 <140> 0\$/09/857,524 <141> 2002-06-21 <150> 60/110,784 <151> 1998 12-03 <160> 17 <170> Microsoft Office 97 <210> 1 <211> 1471 <212> DNA <213> Picramnia pentandra <220> <221> unsure <222> (1402) <223> n = A, C, G, or **T** <400> 1 cttccttgtt cctggaattt tcaaatcact tcctctgttg cacttcaatg gaagagccaa 60 agaagcacat ttcgcaagca gacc \mathbf{t} tgcaa agcataagca accaggagat ttatggatct 120 180 ctatcaaggg aaaagtttac gatat&tcca agtggactaa agagcatccc ggtggtgagc tcccattgtt aagttttgcc ggccaagatg tcactgatgc gttcattgct taccatcctg 240 gcactgcttg gcaatacctt gacaggt ct ttactgggta ctacgttcaa gattactctg 300 tctctgagat gtccaaggac tacagaaggc tcgtctctga gttttctaag atgggtttgt 360 tcaagacacc aggcaaaggg gtctactgot caatcttttt cgtgtctgtg ttgttcgctc 420 tgagtgttta cggtgttctc tactgcaaga gcacctgggc tcatctttgc tctggtttgc taatgggtat gctatggctc cagagtggtt gggtggggca tgattcttgt cactaccaag 480 540 ttatgcctaa ccgtaagctt aatcgtcttt atcaaatcat tgcaggaaat gtgattgctg 600 gtgttagtgt tgcatggtgg aagttggacc ataacaccca tcactttgcc tgtaatagcg 660 ccaatctgga teetgatatt cageacette etataattge catateecca aaatttttea 720 actecettae ateataetat cacaactgea aaatgaeeta tgategeget geeaggtttt 780 ttgttagctt tcagcactgg acattttatc ctgdattgtt aagcgttagg ctctatcttt 840 900 ttattctgtc ttttaaggtg gtgttttcca acaa Δ aaaag ggtatacaag agaagtcagg aaattttagg ctatgcagct ttcttgactt ggtattctct actcctttct cgcctaccca 960 attggcctga aagggtcatg tatttcacgt cctgtt agc agtcgccggg ttccaacatt 1020 ggcagttcag cttgaatcac tttgcttcta atgtttacac tggtttgcct agcggtaatg 1080 attggtttca ccagcagaca aagggcacgc tcaacataac agcttctgct tggtgggatt 1140 ggtttcatgg tggcctgcac tttcagattg agcatcatot gtttccaagg atgcctaagt 1200 gccatttcag gaaaatctca cccattgtga acaaactttg ccagaagcat aatttgtcct 1260 atgaaactgc taccatgtgg gaggccaata aaatggtata ctccaccctg cgtgctgtgg 1320 ctatggaagc taaggatgtt accaagccag ttcccaagaa datggtctgg gaagcaatga 1380 acactttcgg gtgaacctta tnaaacatca agtgctgtct tacccgtaaa agcttccagt 1440 1471 cccaatgttt ctttttttt tttttttt t <210> 2 <211> 448 <212> PRT <213> Picramnia pentandra <400> 2

Ĵ

Met Glu Glu Pro Lys Lys His Ile Ser Gln Ala Asp Leu Ala Lys His Lys Gln Pro Gly Asp Leu Trp Ile Ser Ile Lys Gly Lys Val Tyr Asp Ile Ser Lys Trp Thr Lys Glu His Pro Gly Gly Glu Leu Pro Leu Leu Ser Phe Ala Gly Gln Asp Val Thr Asp Ala Phe Ile Ala Tyr His Pro Gly Thr Ala Trp Gln Tyr Leu Asp Arg Phe Phe Thr Gly Tyr Tyr Val Gln Asp Tyr Ser Val Ser Glu Met Ser Lys Asp Tyr Arg Arg Leu Val Ser Glu Phe Ser Lys Met Gly Leu Phe Lys Thr Pro Gly Lys Gly Val Tyr Cys Ser Ile Phe Phe Val Ser Val Leu Phe Ala Leu Ser Val Tyr Gly Val Leu Tyr Cys Lys Ser Thr Trp Ala His Leu Cys Ser Gly Leu Leu Met Gly Met Leu Trp Leu Gln Ser Gly Trp Val Gly His Asp Ser Cys His Tyr Gln Val Met Pro Asn Arg Lys Leu Asn Arg Leu Phe Gln Ile Ile Ala Gly Asn Val Ile Ala Gly Val Ser Val Ala Trp Trp Lys Leu Asp His Asn Thr His His Phe Ala Cys Asn Ser Ala Asn Leu Asp Pro Asp Ile Gln His Leu Pro Ile Ile Ala Ile Ser Pro Lys Phe Phe Asn Ser Leu Thr Ser Tyr Tyr His Asn Cys Lys Met Thr Tyr Asp Arg Ala Ala Arg Phe Phe Val Ser Phe Gln His Trp Thr Phe Tyr Pro Ala Leu Leu Ser Val Arg Leu Tyr Leu Phe Ile Leu Ser Phe Lys Val Val Phe Ser Asn Asn Lys Arg Val Tyr Lys Arg Ser Gln Glu Ile Leu Gly Tyr Ala Ala Phe Leu Thr Trp Tyr Ser Leu Leu Ser Arg Leu Pro Asn Trp Pro Glu Arg Val Met Tyr Phe Thr Ser Cys Leu Ala Val Ala Gly Phe Gln His Trp Gln Phe Ser Leu Asn His Phe Ala Ser Asn Val

325 335 330 Tyr Thr Gly Leu Pro Ser Gly Asn Asp Trp Phe His Gln Gln Thr Lys 350 340 345 Gly Thr Leu Asn Ile Thr Ala Ser Ala Trp Trp Asp Trp Phe His Gly 355 360 365 Gly Leu His Phe Gln Ile Glu His His Leu Phe Pro Arg Met Pro Lys 370 375 380 Cys His Phe Arg Lys Ile Ser Pro Ile Val Asn Lys Leu Cys Gln Lys 395 385 390 400 His Asn Leu Ser Tyr Glu Thr Ala Thr Met Trp Glu Ala Asn Lys Met 405 410 415 Val Tyr Ser Thr Leu Arg Ala Val Ala Met Glu Ala Lys Asp Val Thr 425 430 420 Lys Pro Val Pro Lys Asn Met Val Trp Glu Ala Met Asn Thr Phe Gly 440 445 435 <210> 3 <211> 1764 <212> DNA <213> Zea mays <400> 3 gcacgagete ectetetete eccaateete eccgeeteee ectaecaaat cageaceace caaggegeat cegageeaeg geegegeaat geegeeetet gtegatgeaa tgeeggeeee 120 cggcgacgcc gcgggcgccg gcgacgtgcg catgatetee tecaaggage tecgegetea 180 cgcttccgcc gacgacctct ggatctccat ctccggcgac gtgtacgacg tcacgccctg 240 gctcccccac caccogggcg gcgacctccc gcttctcacc ctggcggggc aggacgccac 300 cgacgcette geogeetace accegeeete ggegegeeeg etecteegee gettettegt 360 tggccgcctc tctgactacg ccgtctcccc cgcgtccgcc gactaccgcc gcctcctcgc 420 gcagctatec teegegggee tettegaaeg egteggeeee acceesaagg teeagetegt 480 cctgatggcc gtcctcttct acgccgcgct gtacctcgtc ctcgcatgcg ccagcgcctg 540 600 ggcgcacctc ctcgcggggg gtctcattgg cttcgtctgg atccagtccg gctggatggg ccacgacteg ggecaceace geateacegg ceateeggte etegacegeg tegtgeaggt 660 720 gctctccggg aactgcctca ccggcctcag catcgcctgg tggaagtgta accacaacac 780 gcaccacate geetgeaaca geetggaeea tgaeeeggae etceageaea tgeegetett tgccgtctcc cccaagctgt tcggcaacat atggtcctac ttctaccaac ggaccctggc 840 gttcgatgcc gcctcgaaat tcttcatcag ctaccagcac tggaccttct acccggtaat 900 gtgcatcgcc aggataaatc ttctcgcgca gtccgccctg ttcgttctca cggagaagag 960 ggtgccgcag cggttgcttg agatcgcggg ggtcgccaca ttctgggctt ggtacccgtt 1020 gctggtggct tccctgccga attggtggga gagggtcgcg tttgtgcttt tcagcttcac 1080 catctgcggg attcagcacg tccaattctg cctgaaccac ttctcgtccg acgtgtatgt 1140 cgggccaccc aagggcaatg actggtttga gaagcagacg gcaggcacgc tcgacatcct 1200 gtgctctcct tggatggatt ggttccacgg tggcctgcag ttccagattg agcaccatct 1260 gtttccccgc ctacctcggt gccaccttcg caaggttgca ccggccgtcc gcgacctttg 1320 caagaagcat gggctcactt attctgcagc cacattctgg ggtgcaaatg tgcttacatg 1380 gaagacactc agggctgctg cattgcaggc caggaccgct acaagtggtg gtgctccgaa 1440 gaatttggta tgggaggctg tgaacaccca tggataaatg ggatgaagat acgggctaat 1500 ggcaacttct ggtgttcagc ttggtgccca tgtgattgtc tggatgcctt tcagttattt 1560 agagatattq atcattcaac ctgcctgagt caggttggaa ttttcgtgtt gacaagtggc 1620 tgtctatcca gttggagagt tcatgcttca atagtctggt tgttcacggg atgttctgtt 1680 ctccctatca cggtaactat atgatgatga tccttgcttt aattcatgaa cacttgtttc 1740 1764 aagattaaaa aaaaaaaaaa aaaa

60

<210> 4

<211> 462
<212> PRT
<213> Zea mays

<400> 4 Met Pro Pro Ser Val Asp Ala Met Pro Ala Pro Gly Asp Ala Ala Gly - 5 Ala Gly Asp Val Arg Met Ile Ser Ser Lys Glu Leu Arg Ala His Ala Ser Ala Asp Asp Leu Trp Ile Ser Ile Ser Gly Asp Val Tyr Asp Val Thr Pro Trp Leu Pro His His Pro Gly Gly Asp Leu Pro Leu Leu Thr Leu Ala Gly Gln Asp Ala Thr Asp Ala Phe Ala Ala Tyr His Pro Pro Ser Ala Arg Pro Leu Leu Arg Arg Phe Phe Val Gly Arg Leu Ser Asp Tyr Ala Val Ser Pro Ala Ser Ala Asp Tyr Arg Arg Leu Leu Ala Gln Leu Ser Ser Ala Gly Leu Phe Glu Arg Val Gly Pro Thr Pro Lys Val Gln Leu Val Leu Met Ala Val Leu Phe Tyr Ala Ala Leu Tyr Leu Val Leu Ala Cys Ala Ser Ala Trp Ala His Leu Leu Ala Gly Gly Leu Ile Gly Phe Val Trp Ile Gln Ser Gly Trp Met Gly His Asp Ser Gly His His Arg Ile Thr Gly His Pro Val Leu Asp Arg Val Val Gln Val Leu Ser Gly Asn Cys Leu Thr Gly Leu Ser Ile Ala Trp Trp Lys Cys Asn His Asn Thr His His Ile Ala Cys Asn Ser Leu Asp His Asp Pro Asp Leu Gln His Met Pro Leu Phe Ala Val Ser Pro Lys Leu Phe Gly Asn Ile Trp Ser Tyr Phe Tyr Gln Arg Thr Leu Ala Phe Asp Ala Ala Ser Lys Phe Phe Ile Ser Tyr Gln His Trp Thr Phe Tyr Pro Val Met Cys Ile Ala Arg Ile Asn Leu Leu Ala Gln Ser Ala Leu Phe Val Leu Thr Glu Lys Arg Val Pro Gln Arg Leu Leu Glu Ile Ala Gly Val Ala Thr

Phe Trp Ala Trp Tyr Pro Leu Leu Val Ala Ser Leu Pro Asn Trp Trp 305 310 315 320 Glu Arg Val Ala Phe Val Leu Phe Ser Phe Thr Ile Cys Gly Ile Gln 335 325 330 His Val Gln Phe Cys Leu Asn His Phe Ser Ser Asp Val Tyr Val Gly 345 340 350 Pro Pro Lys Gly Asn Asp Trp Phe Glu Lys Gln Thr Ala Gly Thr Leu 355 360 365 Asp Ile Leu Cys Ser Pro Trp Met Asp Trp Phe His Gly Gly Leu Gln 370 375 380 Phe Gln Ile Glu His His Leu Phe Pro Arg Leu Pro Arg Cys His Leu 395 385 390 400 Arg Lys Val Ala Pro Ala Val Arg Asp Leu Cys Lys Lys His Gly Leu 410 415 405 Thr Tyr Ser Ala Ala Thr Phe Trp Gly Ala Asn Val Leu Thr Trp Lys 420 425 430 Thr Leu Arg Ala Ala Ala Leu Gln Ala Arg Thr Ala Thr Ser Gly Gly 445 435 440 Ala Pro Lys Asn Leu Val Trp Glu Ala Val Asn Thr His Gly 455 460 450 ^{..} <210> 5 <211> 880 <212> DNA <213> Glycine max <220> <221> unsure <222> (496) <223> n = A, C, G, or T <220> <221> unsure <222> (512) <223> n = A, C, G, or T <220> <221> unsure <222> (523) <223> n = A, C, G, or T <220> <221> unsure <222> (532) <223> n = A, C, G, or T <220> <221> unsure <222> (630) <223> n = A, C, G, or T

<220> <221> unsure <222> (700) <223> n = A, C, G, or T <220> <221> unsure <222> (730) <223> n = A, C, G, or T <220> <221> unsure <222> (738) <223> n = A, C, G, or T <220> <221> unsure <222> (761) <223> n = A, C, G, or T <220> <221> unsure <222> (764) <223> n = A, C, G, or T <220> <221> unsure <222> (814) <223> n = A, C, G, or T <220> <221> unsure <222> (822) <223> n = A, C, G, or T <220> <221> unsure <222> (824) <223> n = A, C, G, or T <220> <221> unsure <222> (838) <223> n = A, C, G, or T <220> <221> unsure, <222> (842) <223> n = A, C, G, or T <220> <221> unsure <222> (876) <223> n = A, C, G, or T <400> 5 ccgcctccct cctcctcccc gccttctcca cctcccaccg tctttccgac cacaccgtct 60 ccgccgcctc ctccgactac cgcaagctct tctccgacct ctccgcgctc aacctcttca 120 accgcaaggg ccacacaacc tccatcctcc tctcccttat tctcaccctt tttcctctct 180 ctgtctgcgg cgtcctcttc tccgacagca ctttcgtgca cgtgctttcc gctgcattga 240 taggetttet etggatteag ageggetgga taggeeaega eteeggeeat taeaaegtga 300

tgctcagccg ccgcctcaac cgcgcaattc agattctctc cggcaacatt ctcgccggaa 360 tcagcatcgg ctggtggaag tggaaccaca acgcccacca cattgcatgc aacagcctcg 420 actatgaccc tgatctgcag cacatgccgg tctttgcagt ttcgtcgcgg ttcttcaatt 480 ccataacctc tcattnctat gggaggaagt tngagtttga ttncattgct angttcttga 540 tctgctacca gcactttact ttttacccgg taatgtgtgt tgccagggtc aacttgtatc 600 tgcagacaat tctgctattg ttttcgaggn gaaaagtgca ggatagagct tgaacataat 660 ggggatcett gtgttttgga ettggtteet ettttagtgn ettgeetgee aaattgggee 720 tgataggggn atgtttgngc ttgctagctt tgctgtttgt nccnatccag cacattcagt 780 tctggttgaa tcaccttgct gaaaatttat atgncgggca cnantgggaa tgactggntg 840 anaatcagac aaggggtcat tggatatctc ttgtgnccct 880 <210> 6 <211> 253 <212> PRT <213> Glycine max <220> <221> UNSURE <222> (161) <223> Xaa = ANY AMINO ACID <220> <221> UNSURE <222> (166) <223> Xaa = ANY AMINO ACID <220> <221> UNSURE <222> (170) <223> Xaa = ANY AMINO ACID <220> <221> UNSURE <222> (173) <223> Xaa = ANY AMINO ACID <220> <221> UNSURE <222> (206) <223> Xaa = ANY AMINO ACID <220> <221> UNSURE <222> (252) <223> Xaa = ANY AMINO ACID <400> 6 Leu Pro Ala Phe Ser Thr Ser His Arg Leu Ser Asp His Thr Val Ser 10 15 5 1 Ala Ala Ser Ser Asp Tyr Arg Lys Leu Phe Ser Asp Leu Ser Ala Leu 25 30 20 Asn Leu Phe Asn Arg Lys Gly His Thr Thr Ser Ile Leu Leu Ser Leu 40 45 35 Ile Leu Thr Leu Phe Pro Leu Ser Val Cys Gly Val Leu Phe Ser Asp 50 55 60 Ser Thr Phe Val His Val Leu Ser Ala Ala Leu Ile Gly Phe Leu Trp 65 70 75 80

Ile Gln Ser Gly Trp Ile Gly His Asp Ser Gly His Tyr Asn Val Met 95 85 90 Leu Ser Arg Arg Leu Asn Arg Ala Ile Gln Ile Leu Ser Gly Asn Ile 105 110 100 Leu Ala Gly Ile Ser Ile Gly Trp Trp Lys Trp Asn His Asn Ala His 120 125 115 His Ile Ala Cys Asn Ser Leu Asp Tyr Asp Pro Asp Leu Gln His Met 130 135 140 Pro Val Phe Ala Val Ser Ser Arg Phe Phe Asn Ser Ile Thr Ser His 155 160 145 150 Xaa Tyr Gly Arg Lys Xaa Glu Phe Asp Xaa Ile Ala Xaa Phe Leu Ile 170 175 165 Cys Tyr Gln His Phe Thr Phe Tyr Pro Val Met Cys Val Ala Arg Val 180 185 190 Asn Leu Tyr Leu Gln Thr Ile Leu Leu Leu Phe Ser Arg Xaa Lys Val 195 200 205 Gln Asp Arg Ala Leu Asn Ile Met Gly Ile Leu Val Phe Trp Thr Trp 215 220 210 Phe Leu Phe Leu Leu Ala Leu Leu Phe Val Pro Ile Gln His Ile Gln 240 230 235 225 Phe Trp Leu Asn His Leu Ala Glu Asn Leu Tyr Xaa Gly 250 245 <210> 7 <211> 1934 <212> DNA <213> Glycine max <400> 7 gcacgagcac acaagtaaaa ccttagagag agagagagag agagagagag ggtaaaaggg tattagatcc ttgaaccaga tcaaatcatc aaaatctctg tctatggggt tgtgaaaaca gaagaaccag ataccccatt attgttctta tctatctatc tgtctatatt ctatttatct tottattgta gttotcattg tgtotgattt cagtgatttg tgttgttttt ggttaacaca agcaatggag gttgttgaga aggagaagaa gtacataacc tcagaggagc tgaagggtca caacaaggag ggagatttat ggatctcaat tcaaggtaag gtgtacaatg tctcagattg ggtcaaggag caccetggtg gtgatgttee aateteaaae ettgetggee aggatgteae tgatgcattc atagcatacc atcctggcac agcatggtca caccttgaaa aattcttcac tggctaccac ctcagtgact tcaaggtctc tgaggtgtcc aaagactaca gaaagcttgc atctgagttc tcaaaattgg gtctttttga caccaaaggg catgtcactt catgcaccct tgcatctgtt gctgttatgt tcctcattgt actctatggt gttctgaggt gcactagtgt gtgggctcat ttgggttcag gcatgctctt agggttgctt tggatgcaaa gtgcttatgt gggccatgat tctggccact atgtggttat gacaaccaat ggtttcaaca aggttgcaca gatectetet gggaactget tgacegggat aageattget tggtggaagt ggaeteacaa tgctcaccac attgcgtgca acagccttga ccatgaccct gatctgcagc acatgccggt ctttgcagtt tcgtcgcggt tcttcaattc cataacctct catttctatg ggaggaagtt 1020 ggagtttgat ttcattgcta ggttcttgat ctgctaccag cactttactt tttacccggt 1080 aatgtgtgtt gccagggtca acttgtatct gcagacaatt ctgctattgt tttcgaggcg 1140 aaaagtgcag gatagagcct tgaacataat ggggatcctt gtgttttgga cttggttccc 1200 tcttttagtg tcttgcctgc caaattggcc tgagagggtt atgtttgtgc ttgctagctt 1260

60

120

180

240

300

360

420

480

540

600

660 720

780

840

900

tgctgtttgt tccatccage acattcagtt ctgtttgaat cactttgctg caaatgtata 1320 tgtcqggcca ccgagtggga atgactggtt tgagaagcag acaagtggta cattggatat 1380 ctcttgtgcc tcttcgatgg attggttttt cggtggcttg cagtttcagc ttgagcatca 1440 tttgtttcca aggctacctc ggtgccaatt gaggaagatt tcgcctttgg ttagtgacct 1500 ttgcaagaag cataatttgc cttataggag cttgtcattt tgggaggcca atcagtggac 1560 aattaggacc ctcaggactg ctgccctaca agctagggac ttaacaaacc ctgcccctaa 1620 gaatttgttg tgggaagctg ttaataccca tggctgaggc atttggagtt tttagagttt 1680 aggattttgt caaggtcttt ttttttttg ttttctcttt aaaaagaaaa aaaattctca 1740 ttgtgatttt gctagccccc acttttccag attgggcttt gaatttaact ttttgttagg 1800 tgtggtgtac aaatggatgg tgatccagat gttactgcag ttcatgtgct ttgcatcaat 1860 aaaaaaaaa aaaa 1934 <210> 8 <211> 450 <212> PRT <213> Glycine max <400> 8 Met Glu Val Val Glu Lys Glu Lys Lys Tyr Ile Thr Ser Glu Glu Leu 10 1 Lys Gly His Asn Lys Glu Gly Asp Leu Trp Ile Ser Ile Gln Gly Lys 20 25 Val Tyr Asn Val Ser Asp Trp Val Lys Glu His Pro Gly Gly Asp Val 40 45 35 Pro Ile Ser Asn Leu Ala Gly Gln Asp Val Thr Asp Ala Phe Ile Ala 50 55 Tyr His Pro Gly Thr Ala Trp Ser His Leu Glu Lys Phe Phe Thr Gly 75 70 65 Tyr His Leu Ser Asp Phe Lys Val Ser Glu Val Ser Lys Asp Tyr Arg 90 85 Lys Leu Ala Ser Glu Phe Ser Lys Leu Gly Leu Phe Asp Thr Lys Gly 105 110 100 His Val Thr Ser Cys Thr Leu Ala Ser Val Ala Val Met Phe Leu Ile 125 115 120 Val Leu Tyr Gly Val Leu Arg Cys Thr Ser Val Trp Ala His Leu Gly 135 140 130 Ser Gly Met Leu Leu Gly Leu Leu Trp Met Gln Ser Ala Tyr Val Gly 160 155 150 145 His Asp Ser Gly His Tyr Val Val Met Thr Thr Asn Gly Phe Asn Lys 170 175 165 Val Ala Gln Ile Leu Ser Gly Asn Cys Leu Thr Gly Ile Ser Ile Ala 190 180 185 Trp Trp Lys Trp Thr His Asn Ala His His Ile Ala Cys Asn Ser Leu 205 195 200 Asp His Asp Pro Asp Leu Gln His Met Pro Val Phe Ala Val Ser Ser 220 215 210

Arg Phe Phe Asn Ser Ile Thr Ser His Phe Tyr Gly Arg Lys Leu Glu Phe Asp Phe Ile Ala Arg Phe Leu Ile Cys Tyr Gln His Phe Thr Phe Tyr Pro Val Met Cys Val Ala Arg Val Asn Leu Tyr Leu Gln Thr Ile Leu Leu Phe Ser Arg Arg Lys Val Gln Asp Arg Ala Leu Asn Ile Met Gly Ile Leu Val Phe Trp Thr Trp Phe Pro Leu Leu Val Ser Cys Leu Pro Asn Trp Pro Glu Arg Val Met Phe Val Leu Ala Ser Phe Ala Val Cys Ser Ile Gln His Ile Gln Phe Cys Leu Asn His Phe Ala Ala Asn Val Tyr Val Gly Pro Pro Ser Gly Asn Asp Trp Phe Glu Lys Gln Thr Ser Gly Thr Leu Asp Ile Ser Cys Ala Ser Ser Met Asp Trp Phe Phe Gly Gly Leu Gln Phe Gln Leu Glu His His Leu Phe Pro Arg Leu Pro Arg Cys Gln Leu Arg Lys Ile Ser Pro Leu Val Ser Asp Leu Cys Lys Lys His Asn Leu Pro Tyr Arg Ser Leu Ser Phe Trp Glu Ala Asn Gln Trp Thr Ile Arg Thr Leu Arg Thr Ala Ala Leu Gln Ala Arg Asp Leu Thr Asn Pro Ala Pro Lys Asn Leu Leu Trp Glu Ala Val Asn Thr His Gly <210> 9 <211> 1972 <212> DNA <213> Triticum aestivum <400> 9 gcacgagete ectaacaaae eteegttget gttttaagat eegateteee etteeeeet cccctccctt cctcctgagt cctgaccacc cctcctcgcg ctccagctaa atccacgcca 120 ccgatggccc gcacgggcct cgcggacgca acggcgccgg aagccgacgc aatgccggcc 180 gccaqcaaqq acqccgccga cgtccgcatg atetecacca aggagetgca ggcgcacget 240 geogeggaeg acctetggat etecatetee ggggaegtet acgaegteae geogtggetg 300 cgccaccacc cgggcggcga ggtcccgctc atcaccctcg ccggccagga cgccaccgac 360 gccttcatgg cctaccaccc gccctccgtg cgcccgctcc tccgccgctt cttcgtcggc 420 cgcctctccg actacaccgt cccccccgcc tccgccgact tccgccgcct cctcgcgcag ctctcctccg cgggcctctt cgagcgcgtc ggccacaccc ccaagttcct gctcgtcgca 540 atgteggtge tettetgeat egecetetae tgegteeteg eetgeteeag eaceggggee 600

cacatgttcg ccggcggcct cattggcttc atctggatcc agtcgggctg gattggccat 660 gactccggcc accaccaaat caccaggcac cccgcgctca accgcctcct gcaggtggtc 720 teegggaaet geeteacegg eeteggeate geetggtgga agtteaaeea caacaeaeae 780 cacateteet geaacageet egaceatgae eeggacetee ageaettgee getettegeg 840 gtttccacca agetettcaa caacetttgg teggtetget acgagegeae ettggegttt 900 gatgccatat ccaagttctt cgtcagctac cagcactgga cattctaccc ggtgatggga 960 tttgcaagga taaatcttct tgtgcagtca atcgtgttcc tgatcacgca aaagaaggtg 1020 cggcagcgtt ggctggagat cgccggagtt gcagcgttct gggtttggta ccccttgctg 1080 gtctcttgcc tgccgaattg gtgggagagg gttgcttttg tgcttgcaag ctttgtgatc 1140 acggggattc agcatgttca gttctgcctg aaccacttct catccgctgt gtatgttggg 1200 ccaccaaagg ggaacgactg gtttgagagg caaacagcgg gcacacttga tatcaagtgc 1260 tccccgtgga tggattggtt ccatggtggt ctgcagttcc aggttgaaca ccatttgttt 1320 cctcgcctgc ctcgctgcca ctataggatg gtcgcgccga ttgtgcgtga cctttgcaag 1380 aagcatgggc tgtcttatgg tgccgccaca ttctgggagg caaatgtaat gacatggaag 1440 acgctaaggg ctgcagcatt gcaggccagg gaagccacta ctggagctgc tccaaagaat 1500 ctggtctggg aagctttgaa cactcatgga tgactgggat caggactgga gtatgagaca 1560 attgtaagcg tcgagccttg cgtgcatgca gttatctgat tgcttctcga ttgcgtagag 1620 atattgatcc ttttagctgt tggaatcgtg ttggattttt cgtgttgcca ggtgactatc 1680 tttgcagttc aatcgtgggt tcatgcttca gttgtgtact tgtacaccat atttagattg 1740 ttgggttctc cctatcatgg taactacatc aatagtactt gatttacatc ataaaatccg 1800 tggettatet ttacatecat tteattttge ttgeaagtte atgaaaetgt aaaeteaatt 1860 gatggtttgt agcgtgtata tcctgctgct atggcagctt gaactgcatt ttgggaacat 1920 1972 <210> 10 <211> 469 <212> PRT <213> Triticum aestivum <400> 10 Met Ala Arg Thr Gly Leu Ala Asp Ala Thr Ala Pro Glu Ala Asp Ala 10 1 15 Met Pro Ala Ala Ser Lys Asp Ala Ala Asp Val Arg Met Ile Ser Thr 20 25 30 Lys Glu Leu Gln Ala His Ala Ala Ala Asp Asp Leu Trp Ile Ser Ile 35 40 45 Ser Gly Asp Val Tyr Asp Val Thr Pro Trp Leu Arg His His Pro Gly 55 60 50 Gly Glu Val Pro Leu Ile Thr Leu Ala Gly Gln Asp Ala Thr Asp Ala 65 70 75 80 Phe Met Ala Tyr His Pro Pro Ser Val Arg Pro Leu Leu Arg Arg Phe 90 85 Phe Val Gly Arg Leu Ser Asp Tyr Thr Val Pro Pro Ala Ser Ala Asp 100 105 110 Phe Arg Arg Leu Leu Ala Gln Leu Ser Ser Ala Gly Leu Phe Glu Arg 120 125 115 Val Gly His Thr Pro Lys Phe Leu Leu Val Ala Met Ser Val Leu Phe 130 135 140 Cys Ile Ala Leu Tyr Cys Val Leu Ala Cys Ser Ser Thr Gly Ala His 155 160 145 150 Met Phe Ala Gly Gly Leu Ile Gly Phe Ile Trp Ile Gln Ser Gly Trp

Ile Gly His Asp Ser Gly His His Gln Ile Thr Arg His Pro Ala Leu Asn Arg Leu Leu Gln Val Val Ser Gly Asn Cys Leu Thr Gly Leu Gly Ile Ala Trp Trp Lys Phe Asn His Asn Thr His His Ile Ser Cys Asn Ser Leu Asp His Asp Pro Asp Leu Gln His Leu Pro Leu Phe Ala Val Ser Thr Lys Leu Phe Asn Asn Leu Trp Ser Val Cys Tyr Glu Arg Thr Leu Ala Phe Asp Ala Ile Ser Lys Phe Phe Val Ser Tyr Gln His Trp Thr Phe Tyr Pro Val Met Gly Phe Ala Arg Ile Asn Leu Leu Val Gln Ser Ile Val Phe Leu Ile Thr Gln Lys Lys Val Arg Gln Arg Trp Leu Glu Ile Ala Gly Val Ala Ala Phe Trp Val Trp Tyr Pro Leu Leu Val Ser Cys Leu Pro Asn Trp Trp Glu Arg Val Ala Phe Val Leu Ala Ser Phe Val Ile Thr Gly Ile Gln His Val Gln Phe Cys Leu Asn His Phe Ser Ser Ala Val Tyr Val Gly Pro Pro Lys Gly Asn Asp Trp Phe Glu Arg Gln Thr Ala Gly Thr Leu Asp Ile Lys Cys Ser Pro Trp Met Asp Trp Phe His Gly Gly Leu Gln Phe Gln Val Glu His His Leu Phe Pro Arg Leu Pro Arg Cys His Tyr Arg Met Val Ala Pro Ile Val Arg Asp Leu Cys Lys Lys His Gly Leu Ser Tyr Gly Ala Ala Thr Phe Trp Glu Ala Asn Val Met Thr Trp Lys Thr Leu Arg Ala Ala Ala Leu Gln Ala Arg Glu Ala Thr Thr Gly Ala Ala Pro Lys Asn Leu Val Trp Glu Ala Leu Asn Thr His Gly <210> 11 <211> 448

165 [•]

<212> PRT <213> Borago officinalis

.

· ·

·

••

· .

<400> 11 Met Ala A 1	la Gln 1	Ile Lys 5	Lys	Tyr	Ile	Thr 10	Ser	Asp	Glu	Leu	Lys 15	Asn
- His Asp L	ys Pro (20	-	Leu	Trp	Ile 25		Ile	Gln	Gly	Lys 30		Tyr
Asp Val S	er Asp 1 35	[rp Val	Lys	Asp 40	His	Pro	Gly	Gly	Ser 45	Phe	Pro	Leu
Lys Ser L 50	eu Ala G	Gly Gln	Glu 55	Val	Thr	Asp	Ala	Phe 60	Val	Ala	Phe	His
Pro Ala S 65	er Thr I	frp Lys 70	Asn	Leu	Asp	Lys	Phe 75	Phe	Thr	Gly	Tyr	Tyr 80
Leu Lys A	sp Tyr S	Ser Val 85	Ser	Glu	Val	Ser 90	Lys	Asp	Tyr	Arg	Lys 95	Leu
Val Phe G	lu Phe S 100	Ser Lys	Met	Gly	Leu 105	Ťyr	Asp	Lys	Lys	Gly 110	His	Ile
Met Phe A 1	la Thr I 15	Leu Cys	Phe	Ile 120	Ala	Met	Leu	Phe	Ala 125	Met	Ser	Val
Tyr Gly V 130	al Leu B	Phe Cys	Glu 135	Gly	Val	Leu	Val	His 140	Leu	Phe	Ser	Gly
Cys Leu M 145	et Gly E	he Leu? 150	Trp	Ile	Gln	Ser	Gly 155	Trp	Ile	Gly	His	Asp 160
Ala Gly H	-	4et Val 165	Val	Ser	Asp	Ser 170	Arg	Leu	Asn	Lys	Phe 175	Met
Gly Ile P	he Ala A 180	Ala Asn	Cys	Leu	Ser 185	Gly	Ile	Ser	Ile	Gly 190	Trp	Trp
Lys Trp A 1	sn His <i>F</i> 95	Asn Ala	His	His 200	Ile	Ala	Cys	Asn	Ser 205	Leu	Glu	Tyr
Asp Pro A 210		Gln Tyr	Ile 215	Pro	Phe	Leu	Val	Val 220	Ser	Ser	Lys	Phe
Phe Gly S 225	er Leu 1	Thr Ser 230	His	Phe	Tyr	Glu	Lys 235	Arg	Leu	Thr	Phe	Asp 240
Ser Leu S		Phe Phe 245	Val	Ser	Tyr	Gln 250	His	Trp	Thr	Phe	Tyr 255	Pro
Ile Met C	ys Ala <i>F</i> 260	Ala Arg	Leu	Asn	Met 265	Tyr	Val	Gln	Ser	Leu 270	Ile	Met
Leu Leu T 2	hr Lys <i>F</i> 75	Arg Asn	Val	Ser 280	Tyr	Arg	Ala	His	Glu 285	Leu	Leu	Gly
Cys Leu V 290	al Phe S	Ser Ile	Trp 295	Tyr	Pro	Leu	Leu	Val 300	Ser	Cys	Leu	Pro

Asn Trp Gly Glu Arg Ile Met Phe Val Ile Ala Ser Leu Ser Val Thr Gly Met Gln Gln Val Gln Phe Ser Leu Asn His Phe Ser Ser Val Tyr Val Gly Lys Pro Lys Gly Asn Asn Trp Phe Glu Lys Gln Thr Asp Gly Thr Leu Asp Ile Ser Cys Pro Pro Trp Met Asp Trp Phe His Gly Gly Leu Gln Phe Gln Ile Glu His His Leu Phe Pro Lys Met Pro Arg Cys Asn Leu Arg Lys Ile Ser Pro Tyr Val Ile Glu Leu Cys Lys Lys His Asn Leu Pro Tyr Asn Tyr Ala Ser Phe Ser Lys Ala Asn Glu Met Thr Leu Arg Thr Leu Arg Asn Thr Ala Leu Gln Ala Arg Asp Ile Thr Lys Pro Leu Pro Lys Asn Leu Val Trp Glu Ala Leu His Thr His Gly <210> 12 <211> 469 <212> PRT <213> Triticum aestivum <400> 12 Met Ala Arg Thr Gly Leu Ala Asp Ala Thr Ala Pro Glu Ala Asp Ala Met Pro Ala Ala Ser Lys Asp Ala Ala Asp Val Arg Met Ile Ser Thr Lys Glu Leu Gln Ala His Ala Ala Ala Asp Asp Leu Trp Ile Ser Ile Ser Gly Asp Val Tyr Asp Val Thr Pro Trp Leu Arg His His Pro Gly Gly Glu Val Pro Leu Ile Thr Leu Ala Gly Gln Asp Ala Thr Asp Ala Phe Met Ala Tyr His Pro Pro Ser Val Arg Pro Leu Leu Arg Arg Phe Phe Val Gly Arg Leu Thr Asp Tyr Thr Val Pro Pro Ala Ser Ala Asp Phe Arg Arg Leu Leu Ala Gln Leu Ser Ser Ala Gly Leu Phe Glu Arg Val Gly His Thr Pro Lys Phe Leu Leu Val Ala Met Ser Val Leu Phe Cys Ile Ala Leu Tyr Cys Val Leu Ala Cys Ser Ser Thr Gly Ala His

145					1 ° 50					155					160
			- 1	~ `		- 1			- 1		- 1	a 1		a)	
Met	Phe	Ala	GIY	GIY 165	Leu	lle	GIY	Phe	11e 170	Trp	ITe	GIn	Ser	Gly 175	Trp
· Ile	Gly	His	Asp 180	Ser	Gly	His	His	Gln 185	Ile	Thr	Arg	His	Pro 190	Ala	Leu
Asn	Arg	Leu 195	Leu	Gln	Val	Val	Ser 200	Gly	Asn	Cys	Leu	Thr 205	Gly	Leu	Gly
Ile	Ala 210	Trp	Trp	Lys	Phe	Asn 215	His	Asn	Thr	His	His 220	Ile	Ser	Cys	Asn
Ser 225	Leu	Asp	His	Asp	Pro 230	Asp	Leu	Gln	His	Leu 235	Pro	Leu	Phe	Ala	Val 240
Ser	Thr	Lys	Leu	Phe 245	Asn	Asn	Leu	Trp	Ser 250	Val	Cys	Tyr	Glu	Arg 255	Thr
Leu	Ala	Phe	Asp 260	Ala	Ile	Ser	Lys	Phe 265	Phe	Val	Ser	Tyr	Gln 270	His	Trp
Thr	Phe	Tyr 275	Pro	Val	Met	Gly	Phe 280	Ala	Arg	Ile	Asn	Leu 285	Leu	Val	Gln
Ser	Ile 290	Val	Phe	Leu	Ile	Thr 295	Gln	Lys	Lys	Val	Arg 300	Gln	Arg	Trp	Leu
. Glu 305	Ile	Ala	Gly	Val	Ala 310	Ala	Phe	Trp	Val	Trp 315	Tyr	Pro	Leu	Leu	Val 320
Ser	Cys	Leu	Pro	Asn 325	Trp	Trp	Glu	Arg	Val 330	Ala	Phe	Val	Leu	Ala 335	Ser
Phe	Val	Ile	Thr 340	Gly	Ile	Gln	His	Val 345	Gln	Phe	Cys	Leu	Asn 350	His	Phe
Ser	Ser	Ala 355	Val	Tyr	Val	Gly	Pro 360	Pro	Lys	Gly	Asn	Asp 365	Trp	Phe	Glu
Arg	Gln 370	Thr	Ala	Gly	Thr	Leu 375	Asp	Ile	Lys	Cys	Ser 380	Pro	Trp	Met	Asp
Trp 385	Phe	His	Gly	Gly	Leu 390	Gln	Phe	Gln	Val	Glu 395	His	His	Leu	Phe	Pro 400
Arg	Leu	Pro	Arg	Cys 405	His	Tyr	Arg	Met	Val 410	Ala	Pro	Ile	Val	Arg 415	Asp
Leu	Cys	-	Lys 420	His	Gly	Leu	Ser	Tyr 425	Gly	Ala	Ala	Thr	Phe 430	Trp	Glu
Ala	Asn	Val 435	Met	Thr	Trp	Lys	Thr 440	Leu	Arg	Ala	Ala	Ala 445	Leu	Gln	Ala
Arg	Glu 450	Ala	Thr	Thr	Gly	Ala 455	Ala	Pro	Lys	Asn	Leu 460	Val	Trp	Glu	Ala
Leu 465	Asn	Thr	His	Gly								ı			

.

•

.

<210> 13 <211> 458 <212> PRT <213> Helianthus annuus <400> 13 Met Val Ser Pro Ser Ile Glu Val Leu Asn Ser Ile Ala Asp Gly Lys -5 Lys Tyr Ile Thr Ser Lys Glu Leu Lys Lys His Asn Asn Pro Asn Asp Leu Trp Ile Ser Ile Leu Gly Lys Val Tyr Asn Val Thr Glu Trp Ala Lys Glu His Pro Gly Gly Asp Ala Pro Leu Ile Asn Leu Ala Gly Gln Asp Val Thr Asp Ala Phe Ile Ala Phe His Pro Gly Thr Ala Trp Lys His Leu Asp Lys Leu Phe Thr Gly Tyr His Leu Lys Asp Tyr Gln Val Ser Asp Ile Ser Arg Asp Tyr Arg Lys Leu Ala Ser Glu Phe Ala Lys Ala Gly Met Phe Glu Lys Lys Gly His Gly Val Ile Tyr Ser Leu Cys Phe Val Ser Leu Leu Leu Ser Ala Cys Val Tyr Gly Val Leu Tyr Ser Gly Ser Phe Trp Ile His Met Leu Ser Gly Ala Ile Leu Gly Leu Ala Trp Met Gln Ile Ala Tyr Leu Gly His Asp Ala Gly His Tyr Gln Met Met Ala Thr Arg Gly Trp Asn Lys Phe Ala Gly Ile Phe Ile Gly Asn Cys Ile Thr Gly Ile Ser Ile Ala Trp Trp Lys Trp Thr His Asn Ala His His Ile Ala Cys Asn Ser Leu Asp Tyr Asp Pro Asp Leu Gln His Leu Pro Met Leu Ala Val Ser Ser Lys Leu Phe Asn Ser Ile Thr Ser Val Phe Tyr Gly Arg Gln Leu Thr Phe Asp Pro Leu Ala Arg Phe Phe Val Ser Tyr Gln His Tyr Leu Tyr Tyr Pro Ile Met Cys Val Ala Arg Val Asn Leu Tyr Leu Gln Thr Ile Leu Leu Leu Ile Ser Lys Arg Lys

Ile Pro Asp Arg Gly Leu Asn Ile Leu Gly Thr Leu Ile Phe Trp Thr 290 295 300 Trp Phe Pro Leu Leu Val Ser Arg Leu Pro Asn Trp Pro Glu Arg Val 310 315 320 305 Ala Phe Val Leu Val Ser Phe Cys Val Thr Gly Ile Gln His Ile Gln 325 330 335 Phe Thr Leu Asn His Phe Ser Gly Asp Val Tyr Val Gly Pro Pro Lys 340 345 Gly Asp Asn Trp Phe Glu Lys Gln Thr Arg Gly Thr Ile Asp Ile Ala 355 360 365 Cys Ser Ser Trp Met Asp Trp Phe Phe Gly Gly Leu Gln Phe Gln Leu 370 375 380 Glu His His Leu Phe Pro Arg Leu Pro Arg Cys His Leu Arg Ser Ile 390 395 400 385 Ser Pro Ile Cys Arg Glu Leu Cys Lys Lys Tyr Asn Leu Pro Tyr Val 405 410 415 Ser Leu Ser Phe Tyr Asp Ala Asn Val Thr Thr Leu Lys Thr Leu Arg 425 430 420 Thr Ala Ala Leu Gln Ala Arg Asp Leu Thr Asn Pro Ala Pro Gln Asn 440 445 435 Leu Ala Trp Glu Ala Phe Asn Thr His Gly 455 450 <210> 14 <211> 35 <212> DNA <213> Artificial Sequence <220> <223> Definition of Artificial Sequence: PCR primer for 5' of pk0011.d5 <400> 14 35 tttgcggccg caaatcaatg gaagaagcaa agaag <210> 15 <211> 33 <212> DNA <213> Artificial Sequence <220> <223> Definition of Sequence: antisense PCR primer for 3' of pk0011.d5 <400> 15 33 tttgcggccg ccaggattca cccgaaagtg ttc <210> 16 <211> 823 <212> DNA <213> Triticum aestivum <220>

•

<221> unsure <222> (48) \sim <223> n = A, C, G, or T <220> • <221> unsure <222> (538) <223> n = A, C, G, or T <220> <221> unsure <222> (686) <223> n = A, C, G, or T <220> <221> unsure <222> (704) <223> n = A, C, G, or T <220> <221> unsure <222> (717) <223> n = A, C, G, or T <220> <221> unsure · <222> (727) <223> n = A, C, G, or T <220> <221> unsure <222> (729) <223> n = A, C, G, or T <220> <221> unsure <222> (737) <223> n = A, C, G, or T <220> <221> unsure <222> (741) <223> n = A, C, G, or T <220> <221> unsure <222> (750)..(751) <223> n = A, C, G, or T <220> ` <221> unsure <222> (769) <223> n = A, C, G, or T <220> <221> unsure <222> (777) <223> n = A, C, G, or T <220> <221> unsure

. **^**

<222> (807) <223> n = A, C, G, or T

<400> 16 ctccctaaca aacctccgtt gctgttttaa gatccgatct ccccttcncc cctcccctcc 60 cttcctcctg agtcctgacc acccctcctc gcgctccagc taaatccacg ccaccgatgg 120 cccgcacggg cttcgcggac gcaacggcgc cggaagccga cgcaatgccg gccgccagca 180 aggacgccgc cgacgtccgc atgatctcca ccaaggagct gcaggcgcac gccgccgcgg 240 acgacetetg gatetecate teeggggaeg tetaegaegt caegeeetgg etgegeeace 300 accogggogg cgaggtocog ctcatcacoc togooggoca ggacgcoaco gacgcottca 360 tggcctacca cccgccctcc gtgcgcccgc tcctccgccg cttcttcgtc ggccgcctca 420 ccgactacac tgtccccccc gcctccgccg acttccgccg cctcctcgcg cagctctcct 480 ccgcgggcct cttcgagcgc gtcggcacac ccccaagttc ctgctcgtcg caaagtcngt 540 gctcttctgc atcggcctct actgctcctc gcctgctcaa caccggggcc acatgttcgc 600 cggggggctca ttggcttatc tggtcagtcg ggctggattg gcatactccg gcacacaaat 660 cacaggcacc tgcctcaacg ctctgnagtg gctcgggaat gctnacygct cggatcnctg 720 gggagtnanc acacaanaca nattetgaan ngteacatae etgaeteana tteegtnteg 780 823 ggtcacaagt ctaaaacttg catcgtnaag acttggttag cat <210> 17 <211> 114 <212> PRT <213> Triticum aestivum <400> 17 Met Pro Ala Ala Ser Lys Asp Ala Ala Asp Val Arg Met Ile Ser Thr 10 15 5 1 Lys Glu Leu Gln Ala His Ala Ala Ala Asp Asp Leu Trp Ile Ser Ile 30 25 20 Ser Gly Asp Val Tyr Asp Val Thr Pro Trp Leu Arg His His Pro Gly 35 40 45 Gly Glu Val Pro Leu Ile Thr Leu Ala Gly Gln Asp Ala Thr Asp Ala 50 55 60 Phe Met Ala Tyr His Pro Pro Ser Val Arg Pro Leu Leu Arg Arg Phe 70 75 80 65 Phe Val Gly Arg Leu Thr Asp Tyr Thr Val Pro Pro Ala Ser Ala Asp 90 95 85 Phe Arg Arg Leu Leu Ala Gln Leu Ser Ser Ala Gly Leu Phe Glu Arg 105 110 100

Val Gly